Amendments to the Claims are reflected in the listing of claims which begins on page 3 of this paper.

Remarks/Arguments in response to the Office Action begins on page 12 of this paper.

## I. AMENDMENT

This listing of claims will replace all prior versions, and listings, of claims in the application:

# **Listing of Claims:**

- 1-52. (canceled)
- 53. (withdrawn) The process according to claim 47, wherein said opioid receptor polypeptide is a truncated opioid receptor polypeptide.
- 54. (withdrawn) The process of claim 53, wherein said truncated opioid receptor polypeptide is a truncated kappa or a delta opioid receptor polypeptide.
- 55. (withdrawn) The process of claim 53, wherein said truncated opioid receptor polypeptide comprises amino acid residues 79 to 380 of a kappa opioid receptor polypeptide.
- 56. (withdrawn) The process according to claim 47, wherein said opioid receptor polypeptide is a mutant opioid receptor polypeptide.
- 57. (withdrawn) The process according to claim 56, wherein said mutant opioid receptor polypeptide is a mORD1 polypeptide having an asparagine at residue 95 instead of an aspartate.

58. (withdrawn) The process according to claim 47, wherein providing said opioid receptor polypeptide is transfecting a host cell with a polynucleotide that encodes an opioid receptor polypeptide to form a transformed cell and maintaining said transformed cell under biological conditions sufficient for expression of said opioid receptor polypeptide.

# 59. (canceled)

- 60. (withdrawn) The process of claim 59, wherein the opioid receptor polypeptide comprises a portion of a kappa opioid receptor polypeptide.
- 61. (withdrawn) The process of claim 60, wherein the opioid receptor polypeptide comprises a portion of the second extracellular loop of the kappa opioid receptor polypeptide.
- 62. (withdrawn) The process of claim 61, wherein the opioid receptor polypeptide comprises a negatively charged region of the second extracellular loop of the kappa opioid receptor.

## 63-67. (canceled)

68. (withdrawn) The process of claim 59, wherein the opioid receptor polypeptide comprises a truncated opioid receptor polypeptide.

- 69. (withdrawn) The process of claim 68, wherein said truncated opioid receptor polypeptide is a truncated kappa opioid receptor polypeptide.
- 70. (withdrawn) The process of claim 69, wherein the truncated opioid receptor polypeptide comprises amino acid residues 79 to 380 of a kappa opioid receptor polypeptide.
- 71. (withdrawn) The process of claim 69, wherein the truncated opioid receptor polypeptide comprises amino acid residues 167 to 228 of a kappa opioid receptor polypeptide.
- 72. (withdrawn) The process of claim 59, wherein the candidate specific kappa opioid receptor agonist is pre-screened determining whether the candidate has a positive charge.
- 73. (withdrawn) The process according to claim 59, wherein providing said opioid receptor polypeptide is transfecting a host cell with a polynucleotide that encodes an opioid receptor polypeptide to form a transformed cell and maintaining said transformed cell under biological conditions sufficient for expression of said opioid receptor polypeptide.
- 74. (withdrawn) A specific kappa opioid receptor agonist isolatable by the process of claim 59.
- 75. (withdrawn) The process according to claim 47, wherein said opioid receptor polypeptide is a delta or kappa opioid receptor polypeptide.

- 76. (withdrawn) The process of claim 75, wherein said polypeptide is a delta opioid receptor polypeptide.
- 77. (withdrawn) The process of claim 76, wherein said delta opioid receptor polypeptide comprises the amino acid residue sequence of SEQ ID NO:4.
- 78. (withdrawn) The process of claim 75, wherein said polypeptide is a kappa opioid receptor polypeptide.
- 79. (withdrawn) The process of claim 78, wherein said kappa opioid receptor polypeptide comprises the amino acid sequence of SEQ ID NO:2.
- 80. (withdrawn) The process of claim 78, wherein said kappa opioid receptor polypeptide comprises the amino acid sequence of SEQ ID NO:12.

#### 81-96. (canceled)

- 97. (currently amended) A process of screening a substance for its ability to specifically bind to an opioid receptor, said process comprising the steps of:
  - a) expressing a recombinant opioid receptor polypeptide encoded by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11;
  - b) contacting said substance with the opioid receptor polypeptide; and
  - c) detecting the ability of whether said substance has an ability to specifically bind to said opioid receptor polypeptide.

- 98. (previously presented) The process of claim 97, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 40 contiguous bases of SEQ ID NO:11.
- 99. (previously presented) The process of claim 98, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 50 contiguous bases of SEQ ID NO:11.
- 100. (previously presented) The process of claim 99, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 75 contiguous bases of SEQ ID NO:11.
- 101. (previously presented) The process of claim 100, wherein said opioid receptor polypeptide is encoded for by a nucleic acid sequence comprising at least 100 contiguous bases of SEQ ID NO:11.
- 102. (previously presented) The process of claim 101, wherein said opioid receptor polypeptide is encoded for by a nucleic acid sequence comprising at least 680 contiguous bases of SEQ ID NO:11.

103-108. (canceled)

- 109. (currently amended) A process of isolating a substance with an ability to act as a specific agonist of a kappa opioid receptor, said process comprising the steps of:
  - a) providing an opioid receptor polypeptide comprising the second extracellular loop comprising the amino acid sequence of residues 111 through 136 of SEQ ID NO:12 and encoded for by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11;
  - b) contacting said opioid receptor polypeptide with a composition comprising said substance;

- c) detecting the ability of whether said substance has an ability to bind to agonize said opioid receptor polypeptide; and
- d) isolating said substance if the ability of said substance has an ability to agonize specifically bind to the opioid receptor polypeptide is detected.

# 110-111. (canceled)

- 112. (previously presented) The process of claim 109, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 75 contiguous bases of SEQ ID NO:11.
- 113. (previously presented) The process of claim 112, wherein said opioid receptor polypeptide is encoded for by a nucleic acid sequence comprising at least 100 contiguous bases of SEQ ID NO:11.
- 114. (previously presented) The process of claim 113, wherein said opioid receptor polypeptide is encoded for by a nucleic acid sequence comprising at least 680 contiguous bases of SEQ ID NO:11.

# 115-122. (canceled)

123. (currently amended) The process of claim 143 113, wherein said opioid receptor polypeptide is a kappa opioid receptor polypeptide encoded for by the polynucleotide of SEQ ID NO: 11.

## 124-136. (canceled)

137. (currently amended) A process of screening a substance for its ability to act as a specific agonist of a kappa opioid receptor comprising:

- a) expressing a chimeric recombinant opioid receptor polypeptide comprising the second extracellular loop comprising the amino acid sequence of residues 111 through 136 of SEQ ID NO:12, wherein said chimeric opioid receptor polypeptide is encoded by a nucleic acid sequence comprising 30 contiguous bases of SEQ ID NO:11;
- b) contacting said substance with the opioid receptor polypeptide; and
- c) detecting whether the substance has an the ability of the substance to agonize specifically bind to the opioid receptor polypeptide.
- 138. (previously presented) The process of claim 137, wherein said nucleic acid sequence comprises 40 contiguous bases of SEQ ID NO:11.
- 139. (previously presented) The process of claim 137, wherein said nucleic acid sequence comprises 55 contiguous bases of SEQ ID NO:11.
- 140. (previously presented) The process of claim 137, wherein said nucleic acid sequence comprises 70 contiguous bases of SEQ ID NO:11.
- 141. (previously presented) The process of claim 137, wherein a portion of the chimeric opioid receptor polypeptide comprises SEQ ID NO:14.
- 142. (previously presented) The process of claim 137, wherein the chimeric opioid receptor polypeptide comprises polypeptide portions of both kappa and delta opioid receptors.
- 143. (previously presented) The process according to claim 97 wherein the opioid receptor polypeptide is a kappa opioid receptor polypeptide comprising SEQ ID NO:12.

- 144. (new) A process of screening a substance for its ability to specifically bind to an opioid receptor, said process comprising the steps of:
  - a) expressing a recombinant opioid receptor polypeptide comprising the second extracellular loop comprising the amino acid sequence of residues 111 through 136 of SEQ ID NO:12 and encoded by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11;
  - b) contacting said substance with the opioid receptor polypeptide; and
  - c) detecting whether said substance has an ability to specifically bind to said opioid receptor polypeptide.
- 145. (new) The process of claim 144, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 40 contiguous bases of SEQ ID NO:11.
- 146. (new) The process of claim 145, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 50 contiguous bases of SEQ ID NO:11.
- 147. (new) The process of claim 146, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 75 contiguous bases of SEQ ID NO:11.
- 148. (new) The process of claim 147, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 100 contiguous bases of SEQ ID NO:11.
- 149. (new) The process of claim 148, wherein said opioid receptor polypeptide is encoded by a nucleic acid sequence comprising at least 680 contiguous bases of SEQ ID NO:11.
- 150. (new) The process of claim 97, wherein said substance is an antibody.
- 151. (new) A process of screening a substance for its ability to specifically bind to a recombinant polypeptide encoded by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11, said process comprising the steps of:

- a) expressing a recombinant polypeptide encoded by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11;
- b) contacting said substance with the recombinant polypeptide encoded by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11; and
- c) detecting whether the substance has an ability to specifically bind to said recombinant polypeptide encoded by a nucleic acid sequence comprising at least 30 contiguous bases of SEQ ID NO:11.
- 152. (new) The process of claim 151, wherein said polypeptide is encoded by a nucleic acid sequence comprising at least 40 contiguous bases of SEQ ID NO:11.
- 153. (new) The process of claim 152, wherein said polypeptide is encoded by a nucleic acid sequence comprising at least 50 contiguous bases of SEQ ID NO:11.
- 154. (new) The process of claim 153, wherein said polypeptide is encoded by a nucleic acid sequence comprising at least 75 contiguous bases of SEQ ID NO:11.
- 155. (new) The process of claim 154, wherein said polypeptide is encoded by a nucleic acid sequence comprising at least 100 contiguous bases of SEQ ID NO:11.
- 156. (new) The process of claim 155, wherein said polypeptide is encoded by a nucleic acid sequence comprising at least 680 contiguous bases of SEQ ID NO:11.